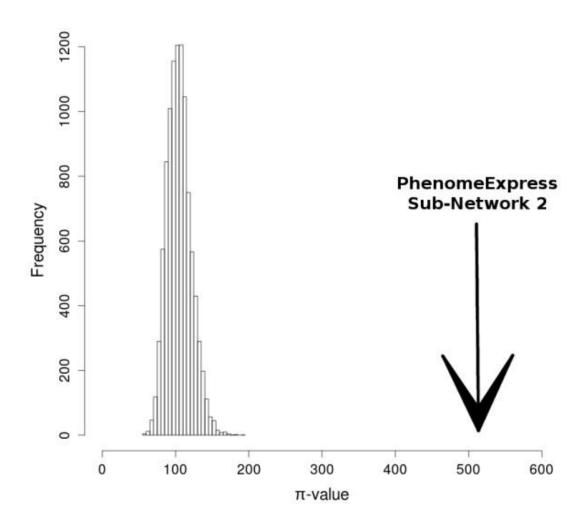
Supplementary Information for:

PhenomeExpress: A refined network analysis of expression datasets by inclusion of known disease phenotypes

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Supplementary Figure S1 – FDR analysis of PhenomeExpress sub-network 2 from the subchondral bone dataset.

Histogram showing the sum of π -values for 10,000 random sub-networks of size equal to the size of sub-network 2 (145 nodes).